

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,817

DATE: 08/29/2002

TIME: 12:21:47

Input Set : A:\SCIOS002C1.TXT

Output Set: N:\CRF3\08292002\J083817.raw

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4 <110> APPLICANT: Schreiner, George F.
        Johnson, Richard J.
 7 <120> TITLE OF INVENTION: METHODS OF TREATING HYPERTENSION AND
        COMPOSITIONS FOR USE THEREIN
11 <130> FILE REFERENCE: SCIOS.002C1
13 <140> CURRENT APPLICATION NUMBER: 10/083,817
14 <141> CURRENT FILING DATE: 2002-02-26
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17 <151> PRIOR FILING DATE: 1998-09-09
19 <150> PRIOR APPLICATION NUMBER: 09/392,932
20 <151> PRIOR FILING DATE: 1999-09-09
22 <160> NUMBER OF SEQ ID NOS: 11
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                                   25
36 Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln
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38 Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu
                           55
40 Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu
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42 Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro
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44 Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His
              100
                                   105
46 Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys
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56 <212> TYPE: PRT
57 <213> ORGANISM: Homo Sapien
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60 Ala Pro Met Ala Glu Gly Gly Gln Asn His His Glu Val Val Lys
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62 Phe Met Asp Val Tyr Gln Arg Ser Tyr Cys His Pro Ile Glu Thr Leu
                              25
64 Val Asp Ile Phe Gln Glu Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys
              40
                                     45
66 Pro Ser Cys Val Pro Leu Met Arg Cys Gly Gly Cys Cys Asn Asp Glu
             55
                                         60
68 Gly Leu Glu Cys Val Pro Thr Glu Glu Ser Asn Ile Thr Met Gln Ile
                                      75
69 65 70
70 Met Arg Ile Lys Pro His Gln Gly Gln His Ile Gly Glu Met Ser Phe
               85
                               90
72 Leu Gln His Asn Lys Cys Glu Cys Arg Pro Lys Lys Asp Arg Ala Arg
73 100
                              105
74 Gln Glu Lys Lys Ser Val Arg Gly Lys Gly Lys Gly Gln Lys Arg Lys
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90 Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly
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                               25
92 Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln
                            40
94 Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu
                       55
96 Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu
                 70
                                      75
98 Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro
                                  90
               85
100 Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His
101 100
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102 Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys
103 115
                            120
104 Glu Cys Arg Pro Lys Lys Asp Arg Ala Arg Gln Glu Asn Pro Cys Gly
                        135
106 Pro Cys Ser Glu Arg Arg Lys His Leu Phe Val Gln Asp Pro Gln Thr
                    150
                                      155
108 Cys Lys Cys Ser Cys Lys Asn Thr Asp Ser Arg Cys Lys Ala Arg Gln
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               165
110 Leu Glu Leu Asn Glu Arg Thr Cys Arg Cys Asp Lys Pro Arg Arg
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168 gtgaagttca tggatgtcta tcagcgcagc tactgccatc caatcgagac cctggtggac 180 169 atcttccagg agtaccctga tgagatcgag tacatcttca agccatcctg tgtgcccctg 240

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Input Set : A:\SCIOS002C1.TXT Output Set: N:\CRF3\08292002\J083817.raw 170 atgcgatgcg ggggctgctg caatgacgag ggcctggagt gtgtgcccac tgaggagtcc 300 171 aacatcacca tgcagattat gcggatcaaa cctcaccaag gccagcacat aggagagatg 360 172 agcttcctac agcacaacaa atgtgaatgc agaccaaaga aagatagagc aagacaagaa 420 173 aaatgtgaca agccgaggcg gtga 175 <210> SEQ ID NO: 7 176 <211> LENGTH: 516 177 <212> TYPE: DNA 178 <213> ORGANISM: Homo Sapien 180 <400> SEQUENCE: 7 181 atgaactttc tgctgtcttg ggtggattgg agccttgcct tgctgctcta cctccaccat 50 182 gecaagtggt cccaggetge acceatggea gaaggaggag ggcagaatea teacgaagtg 120 183 gtgaagttca tggatgtcta tcagcgcagc tactgccatc caatcgagac cctggtggac 180 184 atcttccagg agtaccctga tgagatcgag tacatcttca agccatcctq tqtqcccctq 240 185 atgcgatgcg ggggctgctg caatgacgag ggcctggagt gtgtgcccac tgaggagtcc 300 186 aacatcacca tgcagattat gcggatcaaa cctcaccaag gccagcacat aggagagatg 360 187 agetteetae ageacaacaa atgtgaatge agaccaaaga aagatagage aagacaagaa 420 188 aaaaaatcag ttcgaggaaa gggaaagggg caaaaacgaa agcgcaagaa atcccggtat 480 189 aagteetgga gegtatgtga caageegagg eggtga 516 191 <210> SEQ ID NO: 8 192 <211> LENGTH: 576 193 <212> TYPE: DNA 194 <213> ORGANISM: Homo Sapien 196 <400> SEQUENCE: 8 197 atgaactttc tgctgtcttg ggtgcattgg agcctcgcct tgctgctcta cctccaccat 60 198 gccaagtggt cccaggctgc acccatggca gaaggaggag ggcagaatca tcacgaagtg 120 199 gtgaagttca tggatgtcta tcagcgcagc tactgccatc caatcgagac cctggtggac 180 200 atcttccagg agtaccctga tgagatcgag tacatcttca agccatcctg tgtgccctg 240 201 atgcgatgcg ggggctgctg caatgacgag ggcctggagt gtgtgcccac tgaggagtcc 300 202 aacatcacca tgcagattat gcggatcaaa cctcaccaag gccagcacat aggagagatg 360 203 agetteetae ageacaacaa atgtgaatge agaccaaaga aagatagage aagacaagaa 420 204 aatccctgtg ggccttgctc agagcggaga aagcatttgt ttgtacaaga tccgcagacg 480 205 tgtaaatgtt cctgcaaaaa cacagactcg cgttgcaagg cgaggcagct tgagttaaac 540 206 gaacgtactt gcagatgtga caagccgagg cggtga 576 208 <210> SEO ID NO: 9 209 <211> LENGTH: 642 210 <212> TYPE: DNA 211 <213> ORGANISM: Homo Sapien 213 <400> SEQUENCE: 9 214 atgaactttc tgctgtcttg ggtgcattgg agcctcgcct tgctgctcta cctccaccat 60 215 gccaagtggt cccaggctgc acccatggca gaaggaggag ggcagaatca tcacgaagtg 120 216 gtgaagttca tggatgtcta tcagcgcagc tactgccatc caatcgagac cctggtggac 180 217 atcttccagg agtaccctga tgagatcgag tacatcttca agccatcctg tgtgcccctg 240 218 atgcgatgcg ggggctgctg caatgacgag ggcctggagt gtgtgcccac tgaggagtcc 300 219 aacatcacca tgcagattat gcggatcaaa cctcaccaag gccagcacat aggagagatg 360

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224 ttaaacgaac gtacttgcag atgtgacaag ccgaggcggt ga

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234 gtgaagttca tggatgtcta tcagcgcagc tactgccatc caatcgagac cctggtggac 180
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236 atgegatgeg qqqqctqctq caatqacqaq qqcctqqaqt qtqtqcccac tqaqqaqtcc 300
237 aacatcacca tgcagattat gcggatcaaa cctcaccaag gccagcacat aggagagatg 360
238 agcttcctac agcacaacaa atgtgaatgc agaccaaaga aagatagagc aagacaagaa 420
239 aaaaaatcag ttcgaggaaa gggaaagggg caaaaacgaa agcgcaagaa atcccggtat 480
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241 ccccatccct gtgggccttg ctcagagcgg agaaagcatt tgtttgtaca agatccgcag 600
242 acgtgtaaat gttcctgcaa aaacacagac tcgcgttgca aggcgaggca gcttgagtta 660
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                                40
257 Pro Ser Cys Val Pro Leu Met Arg Cys Gly Gly Cys Cys Asn Asp Glu
                            55
259 Gly Leu Glu Cys Val Pro Thr Glu Glu Ser Asn Ile Thr Met Gln Ile
                        70
261 Met Arg Ile Lys Pro His Gln Gly Gln His Ile Gly Glu Met Ser Phe
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90

105

264

85

100

263 Leu Gln His Asn Lys Cys Glu Cys Arg Pro Lys Lys Asp Arg

VERIFICATION SUMMARYDATE: 08/29/2002PATENT APPLICATION: US/10/083,817TIME: 12:21:48

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